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Form <b>PTO 1449</b>	US Department of Commerce Patent and Trademark Office	ATTY DOCKET NO: P-SH 4810	SERIAL NO: 09/882,359
INFORMATIONAL DISCLOSURE STATEMENT BY APPLICANT		APPLICANT: Shankar Subramaniam	FILING DATE: June 14, 2001
		GROUP: 1645	

### U.S. PATENT DOCUMENTS

EXAM. INITIALS	DOCUMENT NUMBER	DATE	NAME	CLASS	SUB- CLASS	FILING DATE
gaw	5,143,854	09/01/92	Pirrung et al.	17	17	
	5,489,678	02/06/96	Fodor et al.			
	5,571,639	11/05/96	Hubbell et al.			
	5,667,195	09/16/97	McCormick			
	5,744,101	04/28/98	Fodor et al.			

### FOREIGN PATENT DOCUMENTS

EXAM. INITIALS	DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUB- CLASS	TRANSLATION (YES/NO)
gaw	EP1026258A2	08/09/00	Europe			

### OTHER DOCUMENTS (Including Author, Title, Date, Pertinent Pages)

gaw	Adams et al., "The Genome Sequence of <i>Drosophila Melanogaster</i> ," <u>Science</u> , 287:2185-2195 (2000).
gaw	Arsilan et al., "A New Approach to Sequence Comparison: Normalized Sequence Alignment," <u>Bioinformatics</u> , 17(4):327-337 (2001).

EXAMINER <i>gaw</i>	DATE CONSIDERED <i>9/28/02</i>
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EXAMINER: Initial if citation considered, whether or not citation is in conformance with MPEP 609; Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.

Form PTO-419 US Department of Commerce Patent and Trademark Office OIPED 10/10/2001 RECEIVED 100-882,259 100-882,259	ATTY DOCKET NO: P-SH 4810	SERIAL NO: 00882,259
	APPLICANT: Shankar Subramaniam	
INFORMATION DISCLOSURE STATEMENT BY APPLICANT	FILING DATE: June 14, 2001	GROUP: 1645

<i>EW</i>	C. Elegans Sequencing Consortium, "Genome Sequence of the Nematode C. Elegans: A Platform for Investigating Biology," <u>Science</u> , 282:2012-2018 (1998).
	Cheung et al., "Linkage-Disequilibrium Mapping Without Genotyping," <u>Nat. Gen.</u> , 18:225-230 (1998).
	Hayward et al., "Shotgun DNA Microarrays and Stage-Specific Gene Expression in Plasmodium Falciparum Malaria," <u>Mol. Microbiol.</u> , 35(1):6-14 (2000).
	Lin et al., "Whole-Genome Shotgun Optical Mapping of Deinococcus Radiodurans," <u>Science</u> , 285:1558-1562 (1999).
	Mironov et al., "Performance-Guarantee Gene Predictions Via Spliced Alignment," <u>Genomics</u> , 51:332-339 (1998).
	RIKEN Genome Exploration Research Group Phase II Team and FANTOM Consortium, "Functional Annotation of a Full-Length Mouse cDNA Collection," <u>Nature</u> , 409:685-690 (2001).
	Selinger et al., "RNA Expression Analysis Using a 30 Base Pair Resolution Escherichia Coli Genome Array," <u>Nat. Biotech.</u> , 18:1262-1268 (2000).
	Stephan et al., "Positional Cloning Utilizing Genomic DNA Microarrays: The Niemann-Pick Type C Gene as a Model System," <u>Mol. Genet. Metab.</u> , 70:10-18 (2000).
	Sze et al., "Algorithms and Software for Support of Gene Identification Experiments," <u>Bioinformatics</u> , 14(1):14-19 (1998).
	Venter et al., "Shotgun Sequencing of the Human Genome," <u>Science</u> , 280:1540-1542 (1998).
	Venter et al., "The Sequence of the Human Genome," <u>Science</u> , 291:1304-1351 (2001).

EXAMINER <i>EW</i>	DATE CONSIDERED 9/23/02
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